STIC-Biotech/ChemLib

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Chan, Christina

Sent:

Thursday, May 06, 2004 5:54 PM

To:

Graser, Jennifer; STIC-Biotech/ChemLib

Subject: RE: rush search

Please rush. Thanks Chris

Chris Chan

TC 1600 New Hire Training Coordinator and SPE 1644 & 1642 (571)-272-0841

Remsen, 3E89

----Original Message----

From: Graser, Jennifer

Sent: To:

Thursday, May 06, 2004 4:18 PM

Chan, Christina Subject: rush search Importance: High

Hi Christina,

Could you please authorize the following rush search for an application which is due?

Thanks, Jennifer

STIC:

Please search SEQ ID NOs: 12 and 14from 10/625,221 in pending and commercial databases.

Thanks, Jennifer Graser **REMSEN** EO3 B09/3C18 mailbox AU 1645 272-0858

Searcher: Phone: Location: Date Picked Up: ________ Date Completed: Searcher Prep/Review: Clerical:_ Online time:_

TYPE OF SEARCH:	
NA Sequences:	
AA Sequences:	1
Structures:	
Bibliographic:	
Litigation:	
Full text:	•
Patent Family:	
Other:	

VENDOR/COST (where applic.)
STN:	_
DIALOG:	_
Questel/Orbit:	_
DRLink:	
Lexis/Nexis:	-
Sequence Sys.: WWP//C	ZSD
WWW/Internet:	-
Other (specify):	



STIC SEARCH RESUL FEEDBACK FORM

Biotech-Chem Library

Questions about the scope or the results of the search? Contact the searcher or contact:

Mary Hale, Information Branch Supervisor 571-272-2507 Remsen E01 D86

Voluntary Results Feedback Form

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> I am an examiner in Workgroup: Example: 1610
> Relevant prior art found, search results used as follows:
☐ 102 rejection
☐ 103 rejection
Cited as being of interest.
Helped examiner better understand the invention.
Helped examiner better understand the state of the art in their technology.
Types of relevant prior art found:
Foreign Patent(s)
 Non-Patent Literature (journal articles, conference proceedings, new product announcements etc.)
> Relevant prior art not found:
Results verified the lack of relevant prior art (helped determine patentability).
Results were not useful in determining patentability or understanding the invention
Comments:

Drop off or send completed forms to STIC/Biotech-Chem Library Remsen Bldg.



Pending Nucleic Acid and Pending Amino Acid database searches generate two sets of results each. The Pending databases have been split into two parts to reduce the amount of time required for their daily updates. This results in more machine time being available for processing searches.

Searches run against the Nucleic Acid Pending database produce two sets of results, with the extensions rnpm and rnpn

Searches run against the Amino Acid Pending database produce two sets of results, with the extensions rapm and rapn

Because they contain data that is confidential, the results of Pending database searches should not be left in the case.